



PTO/SB/08a/b (08-03)

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Substitute for form 1449A/B/PTO

INFORMATION DISCLOSURE STATEMENT BY APPLICANT

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Sheet	1	of	2	Attorney Docket Number	DU-P02-002
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U.S. PATENT DOCUMENTS

Examiner Initials*	Cite No. ¹	Document Number Number-Kind Code ² (if known)	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear
/JL/	AA	US-6,532,305	03-11-03	LINCOM CORPORATION	
V	AB	US-2004-0083084	04-29-04	WEST	
V	AC	US-2004-0106113	06-03-04	WEST et al.	

FOREIGN PATENT DOCUMENTS

Examiner Initials*	Cite No. ¹	Foreign Patent Document Country Code ³ -Number ⁴ -Kind Code ⁵ (if known)	Publication Date MM-DD-YYYY	Name of Patentee or Applicant of Cited Document	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear	T ⁶

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NON PATENT LITERATURE DOCUMENTS

Examiner Initials	Cite No. ¹	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published.	T ²
/JL/	CA	SELLKE, T. et al., Calibration of pvalues for testing precise null hypotheses, <i>The American Statistician</i> , 55, 62-71, (2001)	
	CB	BREIMAN, L., Statistical Modeling: The two cultures (with discussion), <i>Statistical Science</i> , 16 199-225 (2001)	
	CC	OSBORNE, B.G., Applications of near infrared reflectance spectroscopy to compositional analysis of biscuits and biscuit doughs, <i>J. Sci. Food Agric.</i> , 35, 99-105 (1984)	
	CD	BROWN, P.J., et al., The choice of variables in multivariate regression: A non-conjugate Bayesian decision theory approach, <i>Biometrika</i> , 86, 635-648 (1999).	
	CE	LI, C. et al., Model-based analysis of oligonucleotide arrays: Expression index computation and outlier detection. <i>Proc. Natl. Acad. Sci.</i> , 98, 31-36 (2001)	
	CF	EARLY BREAST CANCER TRIALISTS' COLLABORATIVE GROUP, Polychemotherapy for early breast cancer: an overview of the randomized trials, <i>Lancet</i> , 352:930-942 (2001)	
	CG	WEST, M., et al., Predicting the clinical status of human breast cancer by using gene expression profiles, <i>Proc. Natl. Acad. Sci. USA</i> 98, 11462-11467 (2001)	
	CH	SPANG, R., et al., Prediction and uncertainty in the analysis of gene expression profiles, <i>In Silico Biol.</i> 2, 0033 (2002)	
	CI	VAN T VEER, L.J., et al., Gene expression profiling predicts clinical outcome of breast cancer, <i>Nature</i> 415, 530-536 (2002)	
	CJ	VAN DE VIJVER, M.J., et al., A gene-expression signature as a predictor of survival in breast cancer, <i>N. Engl. J. Med.</i> 347, 1999-2009 (2002)	
	CK	HUANG, E., et al., Gene expression predictors of breast cancer outcomes., <i>Lancet</i> in press (2003)	
	CL	POMEROY, S.L., et al., Prediction of central nervous system embryonal tumour outcome based on gene expression, <i>Nature</i> 415, 436-442 (2002)	
V	CM	ALIZADEH, A.A., et al., Distinct types of diffuse large B-cell lymphoma identified by gene expression profiling, <i>Nature</i> 403, 503-511 (2000)	

Examiner Signature	/Jerry Lin/	Date Considered	09/22/2007
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Substitute for form 1449A/B/PTO				Complete if Known	
				Application Number	10/692002
				Filing Date	October 24, 2003
				First Named Inventor	Mike West
				Art Unit	1631
				Examiner Name	Jerry Lin
Sheet	2	of	2	Attorney Docket Number	DU-P02-002

/JL/	CN	ROSENWALD, A., et al., The use of molecular profiling to predict survival after chemotherapy for diffuse large-B-cell lymphoma.	
	CO	BHATTACHARJEE, A., et al., Classification of human lung carcinomas by mRNA expression profiling reveals distinct adenocarcinoma subclasses, Proc. Natl. Acad. Sci. USA 98, 13790-13795 (2001)	
	CP	RAMASWARMY, S., et al., Multiclass cancer diagnosis using tumor gene expression signatures, Proc. Natl. Acad. Sci. 98, 15149-15154 (2001)	
	CQ	GOLUB, T.R., et al., Molecular classification of cancer: class discovery and class prediction by gene expression monitoring, Science 286, 531-537 (1999)	
	CR	SHIPP, M.A., et al., Diffuse large B-cell lymphoma outcome prediction by gene expression profiling and supervised machine learning, Nat. Med. 8, 68-74 (2002)	
	CS	YEOH, E.-J., et al., Classification, subtype discovery, and prediction of outcome in pediatric acute lymphoblastic leukemia by gene expression profiling, Cancer Cell 1,133-143 (2002)	
	CT	CHENG, et al., Unique Features of Breast Cancer Res. Treat. 2000;63:213-23)	
	CU	MITTRA, I., et al., A Meta-analysis of reported correlations between prognostic factors in breast cancer: does axillary lymph node metastasis represent biology or chronology, Eur.J.Cancer 1991;27:1574-83	
	CV	McGUIRE, W.L., Prognostic factors for recurrence and survival in human breast cancer, Breast Cancer Res Treat. 1987;10:5-9	
	CW	TANDON, A.K., et al., HER-2/neu oncogene protein and prognosis in breast cancer, J.Clin. Oncol. 1989;7:1120-8	
	CX	KASS, R.E., et al., Bayes' factors, J. Am. Stat. Assoc. 90, 773-795 (1998)	
	CY	HOETING, J., et al., Bayesian model averaging, A tutorial; Statistical Science, 14(4), 382-417 (1999)	
	CZ	CLYDE, M., Bayesian Statistics 6, Bernardo J.M. (ed.), pp. 157-185 (Oxford University Press, 1999)	
	CA1	JATOI, I., Significance of axillary lymph node metastasis in primary breast cancer, J. Clin. Oncol. 17, 2334-2340 (1999)	
	CB1	PHILOSOPHOV, L. et al., Medical Diagnostic Decision Rules Based on Mutually Dependent Diagnostic Factors, Comput. Biol. Med., Vol. 27, No. 4, 329-347, 1997.	
	CC1	LIAO, S. et al., Appropriate medical data categorization for data mining classification techniques, Med. Inform. Vol. 27, No. 1, 59-67 (2002).	
	CD1	CHAPMAN, W. et al., A Comparison of Classification Algorithms to Automatically Identify Chest X-Ray Reports That Support Pneumonia, Journal of Biomedical Informatics, 34, 4-14 (2001).	
	CE1	DAWSON, K., et al., A Bayesian approach to the identification of panmictic populations and the assignment of individuals", Genet. Res. Camb., 78, 59-77 (2001).	
	CF1	Copy of International Search Report dated July 1, 2004 from corresponding application no. PCT/US03/33946; citing references PHILOSOPHOV, LIAO, CHAPMAN and DAWSON.	
	CG1	SORLIE, T., et al., Gene expression patterns of breast carcinomas distinguish tumor subclasses with clinical implications, PNAS, Vol. 98, No. 19, 10869-10874 (2001)	
	CH1	D'HAESELEER et al. Bioinformatics, Volume 16, pages 707-726 (2000)	
▼	C1	FRIEDMAN et al., Using Bayesian Networks to Analyze Expression Data, Journal of Computational Biology, Vol. 7, Nos. 3/4, pp. 601-620 (2000).	

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